1/26

SEQUENCE LISTING

<110> Sagami Chemical Research Center

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<151> 1997-11-25

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| | 50 | 55 | 60 | | | |
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| | 65 | 70 | 75 | 80 | | |
| | Cys Val Leu | Val Leu Sei | r Arg Asn | Phe Val G | In Tyr Ala Cys Phe Gly | |
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| | Leu Phe Gly | Ile Ile Ala I | eu Gln Th | r Ile Ala | Tyr Ser Ile Leu Trp | |
| 10 | 100 | 105 | 5 | 110 | | |
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| | 16 | 5 1 | 170 | 175 | | |
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| | Ile Leu Val Ala Ile Gly Phe Lys Thr Lys Leu Ala Ala Leu Thr Leu | | | | | |
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Thr Val Ala Ile Ile Ala Ile Phe Thr Phe Val Glu Pro Val Ser Ala

Val Val Thr Tyr Leu Ile Leu Ala Cys Cys Gln Gly Pro Arg Arg Phe Pro Trp Asn Ile Ile Leu Leu Thr Leu Phe Thr Phe Ala Met Gly Phe Met Thr Gly Thr Ile Ser Ser Met Tyr Gln Thr Lys Ala Val Ile Ile Ala Met Ile Ile Thr Ala Val Val Ser Ile Ser Val Thr Ile Phe Cys Phe Gln Thr Lys Val Asp Phe Thr Ser Cys Thr Gly Leu Phe Cys Val Leu Gly Ile Val Leu Leu Val Thr Gly Ile Val Thr Ser Ile Val Leu Tyr Phe Gln Tyr Val Tyr Trp Leu His Met Leu Tyr Ala Ala Leu Gly Ala Ile Cys Phe Thr Leu Phe Leu Ala Tyr Asp Thr Gln Leu Val Leu Gly Asn Arg Lys His Thr Ile Ser Pro Glu Asp Tyr Ile Thr Gly Ala Leu Gln Ile Tyr Thr Asp Ile Ile Tyr Ile Phe Thr Phe Val Leu Gln Leu Met Gly Asp Arg Asn <210>3 <211>383 <212> PRT

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Cys Gly Pro Gln Cys His Lys Gly Thr Pro Leu Pro Thr Tyr Glu Glu

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Ala Lys Gln Tyr Leu Ser Tyr Glu Thr Leu Tyr Ala Asn Gly Ser Arg

15 85 90 95

Thr Glu Thr Gln Val Gly Ile Tyr Ile Leu Ser Ser Ser Gly Asp Gly

100 105 110

Ala Gln His Arg Asp Ser Gly Ser Ser Gly Lys Ser Arg Arg Lys Arg

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Gln Ile Tyr Gly Tyr Asp Ser Arg Phe Ser Ile Phe Gly Lys Asp Phe

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Leu Leu Asn Tyr Pro Phe Ser Thr Ser Val Lys Leu Ser Thr Gly Cys

145 150 155 160

Thr Gly Thr Leu Val Ala Glu Lys His Val Leu Thr Ala Ala His Cys

25 165 170 175

Ile His Asp Gly Lys Thr Tyr Val Lys Gly Thr Gln Lys Leu Arg Val

180 185 190

Gly Phe Leu Lys Pro Lys Phe Lys Asp Gly Gly Arg Gly Ala Asn Asp Ser Thr Ser Ala Met Pro Glu Gln Met Lys Phe Gln Trp Ile Arg Val Lys Arg Thr His Val Pro Lys Gly Trp Ile Lys Gly Asn Ala Asn Asp Ile Gly Met Asp Tyr Asp Tyr Ala Leu Leu Glu Leu Lys Lys Pro His Lys Arg Lys Phe Met Lys Ile Gly Val Ser Pro Pro Ala Lys Gln Leu Pro Gly Gly Arg Ile His Phe Ser Gly Tyr Asp Asn Asp Arg Pro Gly Asn Leu Val Tyr Arg Phe Cys Asp Val Lys Asp Glu Thr Tyr Asp Leu Leu Tyr Gln Gln Cys Asp Ala Gln Pro Gly Ala Ser Gly Ser Gly Val Tyr Val Arg Met Trp Lys Arg Gln Gln Gln Lys Trp Glu Arg Lys Ile Ile Gly Ile Phe Ser Gly His Gln Trp Val Asp Met Asn Gly Ser Pro Gln Asp Phe Asn Val Ala Val Arg Ile Thr Pro Leu Lys Tyr Ala Gln Ile Cys Tyr Trp Ile Lys Gly Asn Tyr Leu Asp Cys Arg Glu Gly <210>4 <211> 807

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| ggetaccege ageetggeta eggteaceet getggetace cacageceat geeceecace 180 |
|---|
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| tcaggaaagt ctcgaaggaa gcggcagatt tatggctatg acagcaggtt cagcattttt | 420 |
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| gggaaggact teetgeteaa etaceettte teaacateag tgaagttate eaegggetge | 480 |
| accggcaccc tggtggcaga gaagcatgtc ctcacagctg cccactgcat acacgatgg | a 540 |
| aaaacctatg tgaaaggaac ccagaagctt cgagtgggct tcctaaagcc caagtttaaa | 600 |
| gatggtggtc gaggggccaa cgactccact tcagccatgc ccgagcagat gaaatttcag | g 660 |
| tggatccggg tgaaacgcac ccatgtgccc aagggttgga tcaagggcaa tgccaatgac | c 720 |
| ateggeatgg attatgatta tgeceteetg gaacteaaaa ageeecacaa gagaaaattt | 780 |
| atgaagattg gggtgagccc tcctgctaag cagctgccag ggggcagaat tcacttctct | 840 |
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Met Gly Gln Asn Asp

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25 1

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Glu Ser Ser Pro Lys Gln Tyr Met Gln Leu Gly Gly Arg Val Leu Leu

gtt etg atg tte atg ace etc ett eae ttt gae gee age tte ttt tet Val Leu Met Phe Met Thr Leu Leu His Phe Asp Ala Ser Phe Phe Ser att gtc cag aac atc gtg ggc aca gct ctg atg att tta gtg gcc att Ile Val Gln Asn Ile Val Gly Thr Ala Leu Met Ile Leu Val Ala Ile ggt ttt aaa acc aag ctg gct gct ttg act ctt gtt gtg tgg ctc ttt Gly Phe Lys Thr Lys Leu Ala Ala Leu Thr Leu Val Val Trp Leu Phe gee ate aac gta tat tte aac gee tte tgg ace att eea gte tae aag Ala Ile Asn Val Tyr Phe Asn Ala Phe Trp Thr Ile Pro Val Tyr Lys ccc atg cat gac ttc ctg aaa tac gac ttc ttc cag acc atg tcg gtg Pro Met His Asp Phe Leu Lys Tyr Asp Phe Phe Gln Thr Met Ser Val att ggg ggc ttg ctc ctg gtg gtg gcc ctg ggc cct ggg ggt gtc tcc Ile Gly Gly Leu Leu Val Val Ala Leu Gly Pro Gly Gly Val Ser atg gat gag aag aag aag gag tgg taa cagtcacaga tccctacctg Met Asp Glu Lys Lys Lys Glu Trp cctggctaag acccgtggcc gtcaaggact ggttcggggt ggattcaaca aaactgccag 990 ettttatgta teetetteee tteeeteee ttggtaaagg cacagatgtt ttgagaactt 1050 tatttgcaga gacacctgag aatcgatggc tcagtctgct ctggagccac agtctggcgt 1110 ctgaccette agtgeaggee ageetggeag etggaageet eeeceaegee gaggetttgg 1170

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15 25 30 35

Leu Glu Asp Gly Ile Arg Met Trp Phe Gln Trp Ser Glu Gln Arg Asp

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Tyr Ile Asp Thr Trp Asn Cys Gly Tyr Leu Leu Ala Ser Ser Phe

55 60 65

Val Phe Leu Asn Leu Leu Gly Gln Leu Thr Gly Cys Val Leu Val Leu

70 75 80 85

Ser Arg Asn Phe Val Gln Tyr Ala Cys Phe Gly Leu Phe Gly Ile Ile

90 95 100

Ala Leu Gin Thr Ile Ala Tyr Ser Ile Leu Trp Asp Leu Lys Phe Leu

25 105 110 115

Met Arg Asn Leu Ala Leu Gly Gly Gly Leu Leu Leu Leu Ala Glu

120 125 130

Ser Arg Ser Glu Gly Lys Ser Met Phe Ala Gly Val Pro Thr Met Arg Glu Ser Ser Pro Lys Gln Tyr Met Gln Leu Gly Gly Arg Val Leu Leu Val Leu Met Phe Met Thr Leu Leu His Phe Asp Ala Ser Phe Phe Ser Ile Val Gln Asn Ile Val Gly Thr Ala Leu Met Ile Leu Val Ala Ile Gly Phe Lys Thr Lys Leu Ala Ala Leu Thr Leu Val Val Trp Leu Phe Ala Ile Asn Val Tyr Phe Asn Ala Phe Trp Thr Ile Pro Val Tyr Lys Pro Met His Asp Phe Leu Lys Tyr Asp Phe Phe Gln Thr Met Ser Val Ile Gly Gly Leu Leu Val Val Ala Leu Gly Pro Gly Gly Val Ser Met Asp Glu Lys Lys Lys Glu Trp <210>9 <211> 2290 <212> DNA <213> Homo sapiens <400> 9 acacteegag geeaggaaeg eteegtetgg aaeggegeag gteeeageag etggggttee

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Met Ser Asn Pro Ser Ala Pro Pro Pro Tyr Glu

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tee tat get gte tte gtt gte ace tae etg ate ett gee tge tge eag

| | Ser Tyr Ala Val Phe Val Val Thr Tyr Leu Ile Leu Ala Cys Cys Gln | | | | | |
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| | 140 | 145 | 150 | 155 | | |
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| 5 | 160 | 165 | 5 | 170 | | |
| | act ttt gcc atg | ggc ttc atg ac | g ggc acc a | tt tcc agt atg tac caa | 641 | |
| | Thr Phe Ala N | Met Gly Phe M | let Thr Gly | Thr Ile Ser Ser Met | Tyr Gln | |
| | 175 | 180 | 18 | 5 | | |
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| 10 | Thr Lys Ala V | al Ile Ile Ala | Met Ile Ile | Thr Ala Val Val Se | r Ile | |
| | 190 | 195 | 200 | | | |
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| | Ser Val Thr II | e Phe Cys Pho | e Gln Thr L | ys Val Asp Phe Thr | Ser Cys | |
| | 205 | 210 | 215 | | | |
| 15 | aca ggc etc tto | tgt gtc ctg gg | ga att gtg ct | c ctg gtg act ggg att | 785 | |
| | Thr Gly Leu F | he Cys Val L | eu Gly Ile | Val Leu Leu Val Th | r Gly Ile | |
| | 220 | 225 | 230 | 235 | | |
| | | | _ | tac tgg ctc cac atg | 833 | |
| | | - | | yr Val Tyr Trp Leu | His Met | |
| 20 | 240 | 245 | | 250 | | |
| | | | | ctg ttc ctg gct tac | 881 | |
| | - | _ | • | Phe Thr Leu Phe Leu | ı Ala Tyr | |
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| 25 | | | | cac acc atc agc ccc | | |
| 25 | • | | | g Lys His Thr Ile Se | r Pro Glu | |
| | 270 | 275 | 280 | | 077 | |
| | gac tac atc act | ggc gcc ctg c | ag att tac a | ca gac atc atc tac atc | 977 | |

Asp Tyr Ile Thr Gly Ala Leu Gln Ile Tyr Thr Asp Ile Ile Tyr Ile

295 285 290

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Phe Thr Phe Val Leu Gln Leu Met Gly Asp Arg Asn

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25

305

310

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<400> 11

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| | Gly Pro Arg Arg Arg Phe Pro Trp Asn Ile Ile Leu Leu Th | | | eu Thr Leu Phe |
|----|--|-------------------|------------------------|-----------------|
| | 160 | 165 | 170 | |
| | Thr Phe Ala M | let Gly Phe Met | Thr Gly Thr Ile Ser S | Ser Met Tyr Gln |
| | 175 | 180 | 185 | |
| 5 | Thr Lys Ala V | al Ile Ile Ala Me | t Ile Ile Thr Ala Val | Val Ser Ile |
| | 190 | 195 | 200 | |
| | Ser Val Thr Ile | Phe Cys Phe Gl | n Thr Lys Val Asp P | he Thr Ser Cys |
| | 205 | 210 | 215 | |
| | Thr Gly Leu Pl | he Cys Val Leu (| Gly Ile Val Leu Leu | Val Thr Gly Ile |
| 10 | 220 | 225 230 | 235 | |
| | Val Thr Ser Ile | e Val Leu Tyr Pho | e Gln Tyr Val Tyr Ti | rp Leu His Met |
| | 240 | 245 | 250 | |
| | Leu Tyr Ala A | la Leu Gly Ala II | e Cys Phe Thr Leu F | Phe Leu Ala Tyr |
| | 255 | 260 | 265 | |
| 15 | Asp Thr Gln L | eu Val Leu Gly A | Asn Arg Lys His Thr | Ile Ser Pro Glu |
| | 270 | 275 | 280 | |
| | Asp Tyr Ile Th | r Gly Ala Leu G | In Ile Tyr Thr Asp Ile | e Ile Tyr Ile |
| | 285 | 290 | 295 | |
| | Phe Thr Phe V | al Leu Gln Leu N | Met Gly Asp Arg Ası | 1 |
| 20 | 300 | 305 310 |) | |
| | | | | |
| | <210> 11 | | | |
| | <211> 3705 | | · | |
| | <212> DNA | | | |
| 25 | <213> Homo s | apiens | | |
| | | | | |

| | actetegget g | tgcggcggg g | gcaggcatg | g gagccgcgcg cto | ctctcccg gcgc | ccacac | 60 |
|----|---------------|---------------|--|----------------------|---------------|--------|----|
| | ctgtctgagc g | gcgcagcga į | eggg geaggeatgg gageegege eteteteeeg gegeeeaeae 60 gega geegegeee gggegggetg eteggegeg aacagtgete 120 et eea ggg ete ete tte ett ett ett ett ett ett | | | | |
| | ggc atg gca | ggg att cca | ggg ctc ct | e tte ett ete tte tt | t ctg ctc 16 | 58 | |
| | Met Ala | Gly Ile Pro (| Gly Leu L | eu Phe Leu Leu | Phe Phe Leu | Leu | |
| 5 | 1 | 5 | 10 | 15 | | | |
| | tgt gct gtt g | gg caa gtg a | ge eet tae | agt gee eee tgg | aaa ccc act | 216 | |
| | Cys Ala Va | l Gly Gin V | al Ser Pro | Tyr Ser Ala Pro | Trp Lys Pro | Thr | |
| | 2 | 0 | 25 | 30 | | | |
| | tgg cct gca t | tac ege ete e | ct gtc gtc | ttg ccc cag tct ac | ce etc aat 2 | 264 | |
| 10 | Trp Pro Ala | Tyr Arg Le | u Pro Val | Val Leu Pro Gl | n Ser Thr Leu | ı Asn | |
| | 35 | 40 |) | 45 | | | |
| | tta gcc aag | cca gac ttt g | ga gcc gaa | gcc aaa tta gaa | gta tct tct | 312 | |
| | Leu Ala Lys | s Pro Asp Pl | ne Gly Ala | a Glu Ala Lys Lo | eu Glu Val Se | er Ser | |
| | 50 | 55 | (| 50 | | | |
| 15 | tca tgt gga c | ecc cag tgt c | at aag gga | act cca ctg ccc | act tac gaa | 360 | |
| | Ser Cys Gly | Pro Gln Cy | s His Lys | Gly Thr Pro Le | u Pro Thr Tyr | Glu | |
| | 65 | 70 | 75 | | | | |
| | gag gcc aag | caa tat ctg t | ct tat gaa | acg ctc tat gcc a | at ggc agc | 408 | |
| | Glu Ala Lys | s Gln Tyr Le | eu Ser Tyr | Glu Thr Leu Ty | r Ala Asn Gl | y Ser | |
| 20 | 80 | 85 | 90 | 95 | | | |
| | cgc aca gag | acg cag gtg | ggc atc ta | c atc ctc agc agt | agt gga gat | 456 | |
| | Arg Thr Glu | Thr Gln V | al Gly Ile | Tyr Ile Leu Ser | Ser Ser Gly A | Asp | |
| | 10 | 00 | 105 | 110 | | | |
| | ggg gcc caa | cac cga gac | tca ggg to | ct tca gga aag tct | cga agg aag | 504 | |
| 25 | Gly Ala Glr | His Arg A | sp Ser Gly | Ser Ser Gly Ly | s Ser Arg Arg | g Lys | |
| | 115 | 12 | 20 | 125 | | | |
| | caa caa att t | et and tet as | ac age agg | tte age att ttt go | ית פפת תפכ | 552 | |

| | Arg Gln Ile Tyr Gly Tyr Asp Ser Arg Phe Ser Ile Phe Gly Lys Asp | | | | |
|------|---|---|--------------|--|--------------|
| | 130 | 135 | 14 | 40 | |
| | ttc ctg ctc aa | c tac cct ttc t | tca aca tca | gtg aag tta tcc acg gg | gc 600 |
| | Phe Leu Leu | Asn Tyr Pro | Phe Ser | Γhr Ser Val Lys Leu S | Ser Thr Gly |
| 5 | 145 | 150 | 155 | | |
| | tgc acc ggc a | icc ctg gtg go | ca gag aag | cat gtc ctc aca gct gc | c cac 648 |
| • | Cys Thr Gly | Thr Leu Val | l Ala Glu I | Lys His Val Leu Thr | Ala Ala His |
| | 160 | 165 | 170 | 175 | |
| | tgc ata cac g | at gga aaa ac | c tat gtg a | aa gga acc cag aag ct | t cga 696 |
| 10 | Cys Ile His A | Asp Gly Lys | Thr Tyr V | al Lys Gly Thr Gln I | ys Leu Arg |
| | 18 | 0 1 | 185 | 190 | |
| | gtg ggc ttc ci | ta aag ccc aa | g ttt aaa ga | at ggt ggt cga ggg gco | aac 744 |
| | | Leu Lys Pro | Lys Phe I | Lys Asp Gly Gly Arg | Gly Ala Asn |
| | 195 | 200 |) | 205 | |
| 15 | | _ | | atg aaa ttt cag tgg atc | |
| | • | | | In Met Lys Phe Gln | Trp Ile Arg |
| | 210 | 215 | | 20 | |
| | | | | tgg atc aag ggc aat gc | |
| . 20 | vai Lys Arg | 230 | 235 | Gly Trp Ile Lys Gly A | sn Ala Asn |
| . 20 | | | | o ata ann ata ann ana | ccc 888 |
| | | | | c ctg gaa ctc aaa aag Ala Leu Leu Glu Leu | |
| | 240 | 245 | 250 | 255 | Lys Lys I 10 |
| | | | | tg agc cct cct gct aag | cag 936 |
| 25 | | | | ily Val Ser Pro Pro A | |
| _• | 26 | - | 265 | 270 | · |
| | | | | gt tat gac aat gac cga | cca 984 |
| | S 666 E | ,, ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,, | 88 | J | |

Leu Pro Gly Gly Arg Ile His Phe Ser Gly Tyr Asp Asn Asp Arg Pro 275 280 285 ggc aat ttg gtg tat cgc ttc tgt gac gtc aaa gac gag acc tat gac Gly Asn Leu Val Tyr Arg Phe Cys Asp Val Lys Asp Glu Thr Tyr Asp 5 290 295 300 ttg etc tae eag eaa tge gat gee eag eea ggg gee age ggg tet ggg 1080 Leu Leu Tyr Gln Gln Cys Asp Ala Gln Pro Gly Ala Ser Gly Ser Gly 305 310 315 gtc tat gtg agg atg tgg aag aga cag cag cag aag tgg gag cga aaa 1128 10 Val Tyr Val Arg Met Trp Lys Arg Gln Gln Gln Lys Trp Glu Arg Lys 320 325 330 335 att att ggc att ttt tca ggg cac cag tgg gtg gac atg aat ggt tcc Ile Ile Gly Ile Phe Ser Gly His Gln Trp Val Asp Met Asn Gly Ser 340 345 15 cca cag gat ttc aac gtg gct gtc aga atc act cct ctc aaa tat gcc Pro Gln Asp Phe Asn Val Ala Val Arg Ile Thr Pro Leu Lys Tyr Ala 355 360 365 cag att tgc tat tgg att aaa gga aac tac ctg gat tgt agg gag ggg 1272 Gln Ile Cys Tyr Trp Ile Lys Gly Asn Tyr Leu Asp Cys Arg Glu Gly 20 370 375 380 1330 tgacacag tgttccctcc tggcagcaat taagggtctt catgttctta ttttaggaga ggccaaattg ttttttgtca ttggcgtgca cacgtgtgtg tgtgtgtgtg tgtgtaaggt 1390 gtcttataat cttttaccta tttcttacaa ttgcaagatg actggcttta ctatttgaaa 1450 actggtttgt gtatcatatc atatatcatt taagcagttt gaaggcatac ttttgcatag 1510 25 aaataaaaaa aatactgatt tggggcaatg aggaatattt gacaattaag ttaatcttca 1570 cgtttttgca aactttgatt tttatttcat ctgaacttgt ttcaaagatt tatattaaat 1630 atttggcata caagagatat gaattettat atgtgtgcat gtgtgttttc ttctgagatt 1690

10

15

20

25

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aaataattca acatgtccat ctttttagtg ataataaaag aaagcatggt attaaactat 3370
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<212> PRT

<213> Homo sapiens

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1 5 10 15

Cys Ala Val Gly Gln Val Ser Pro Tyr Ser Ala Pro Trp Lys Pro Thr

20 25 30

Trp Pro Ala Tyr Arg Leu Pro Val Val Leu Pro Gln Ser Thr Leu Asn

20 35 40 45

Leu Ala Lys Pro Asp Phe Gly Ala Glu Ala Lys Leu Glu Val Ser Ser

50 55 60

Ser Cys Gly Pro Gln Cys His Lys Gly Thr Pro Leu Pro Thr Tyr Glu

65 70 75

Glu Ala Lys Gln Tyr Leu Ser Tyr Glu Thr Leu Tyr Ala Asn Gly Ser

80 85 90 95

Arg Thr Glu Thr Gln Val Gly Ile Tyr Ile Leu Ser Ser Ser Gly Asp

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| | | 100 | 105 | | 110 | |
|----|-----------|-------------|-----------|-----------|---------------|--------------------|
| | Gly Ala C | in His Arg | Asp Ser | Gly Ser | Ser Gly Lys | Ser Arg Arg Lys |
| | 11 | 5 | 120 | 12 | 5 | |
| | Arg Gln I | le Tyr Gly | Tyr Asp S | Ser Arg I | Phe Ser Ile I | Phe Gly Lys Asp |
| 5 | 130 | 13 | 35 | 140 | | |
| | Phe Leu L | eu Asn Ty | r Pro Phe | Ser Thr | Ser Val Lys | Leu Ser Thr Gly |
| | 145 | 150 | | 155 | | |
| | Cys Thr C | Gly Thr Leu | Val Ala | Glu Lys | His Val Le | u Thr Ala Ala His |
| | 160 | 165 | 1 | 70 | 175 | |
| 10 | Cys Ile H | is Asp Gly | Lys Thr | Γyr Val I | ys Gly Thr | Gln Lys Leu Arg |
| | | 180 | 185 | | 190 | |
| | Val Gly P | he Leu Lys | Pro Lys | Phe Lys | Asp Gly G | y Arg Gly Ala Asn |
| | 19 | 5 | 200 | 20 | 5 | |
| | Asp Ser T | hr Ser Ala | Met Pro | Glu Gln | Met Lys Ph | e Gln Trp Ile Arg |
| 15 | 210 | 2 | 15 | 220 | | |
| | Val Lys A | arg Thr His | Val Pro l | Lys Gly | Trp Ile Lys | Gly Asn Ala Asn |
| | 225 | 230 | | 235 | | |
| | Asp Ile G | ly Met Asp | Tyr Asp | Tyr Ala | Leu Leu Gl | u Leu Lys Lys Pro |
| | 240 | 245 | 2: | 50 | 255 | |
| 20 | His Lys A | arg Lys Phe | Met Lys | Ile Gly | Val Ser Pro | Pro Ala Lys Gln |
| | | 260 | 265 | : | 270 | |
| | Leu Pro C | Gly Gly Arg | Ile His P | he Ser C | Gly Tyr Asp | Asn Asp Arg Pro |
| | 27 | 5 | 280 | 28 | 5 | |
| | Gly Asn I | eu Val Tyı | Arg Phe | Cys Asp | Val Lys A | sp Glu Thr Tyr Asp |
| 25 | 290 | 29 | 95 | 300 | | |
| | Leu Leu 7 | fyr Gln Gln | Cys Asp | Ala Gln | Pro Gly A | a Ser Gly Ser Gly |
| | 305 | 310 | | 315 | | |

